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SEQUENCE LISTING

Shimizu, Nobuyoshi  
Mizuno, Yoshikuni

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<140> 10/776,604

<141> 2004-02-12

<150> 09/601,844

<151> 2000-08-09

<150> PCT/JP99/00545

<151> 1999-02-09

<150> JP 10/27531

<151> 1998-02-09

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| Ser Ile Phe Gln Leu Lys Glu Val Val Ala Lys Arg Gln Gly Val Pro |     |
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| Ala Asp Gln Leu Arg Val Ile Phe Ala Gly Lys Glu Leu Arg Asn Asp |     |
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| tgg act gtg cag aat tgt gac ctg gat cag cag agc att gtt cac att | 308 |
| Trp Thr Val Gln Asn Cys Asp Leu Asp Gln Gln Ser Ile Val His Ile |     |
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| 120 125 130   |     |
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| Ala Gly Ser Pro Ala Gly Arg Ser Ile Tyr Asn Ser Phe Tyr Val Tyr |     |
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| 215 220 225   |     |
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| 230 235 240 245   |     |
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|-----|-----|-----|-----|-----|-----|-----|-----|-------|-----|-----|-----|-----|-----|-----|-----|------|
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| Asp | Cys | Phe | His | Leu | Tyr | Cys | Val | Thr   | Arg | Leu | Asn | Asp | Arg | Gln | Phe |      |
|     |     |     | 265 |     |     |     |     | 270   |     |     |     |     | 275 |     |     |      |
| gtt | cac | gac | cct | caa | ctt | ggc | tac | tcc   | ctg | cct | tgt | gtg | gct | ggc | tgt | 980  |
| Val | His | Asp | Pro | Gln | Leu | Gly | Tyr | Ser   | Leu | Pro | Cys | Val | Ala | Gly | Cys |      |
|     |     | 280 |     |     |     |     | 285 |       |     |     |     | 290 |     |     |     |      |
| ccc | aac | tcc | ttg | att | aaa | gag | ctc | cat   | cac | ttc | agg | att | ctg | gga | gaa | 1028 |
| Pro | Asn | Ser | Leu | Ile | Lys | Glu | Leu | His   | His | Phe | Arg | Ile | Leu | Gly | Glu |      |
|     | 295 |     |     |     |     | 300 |     |       |     |     | 305 |     |     |     |     |      |
| gag | cag | tac | aac | cgg | tac | cag | cag | tat   | ggt | gca | gag | gag | tgt | gtc | ctg | 1076 |
| Glu | Gln | Tyr | Asn | Arg | Tyr | Gln | Gln | Tyr   | Gly | Ala | Glu | Glu | Cys | Val | Leu |      |
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| Gln | Met | Gly | Gly | Val | Leu | Cys | Pro | Arg   | Pro | Gly | Cys | Gly | Ala | Gly | Leu |      |
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| ctg | ccg | gag | cct | gac | cag | agg | aaa | gtc   | acc | tgc | gaa | ggg | ggc | aat | ggc | 1172 |
| Leu | Pro | Glu | Pro | Asp | Gln | Arg | Lys | Val   | Thr | Cys | Glu | Gly | Gly | Asn | Gly |      |
|     |     |     | 345 |     |     |     |     | 350   |     |     |     |     | 355 |     |     |      |
| ctg | ggc | tgt | ggg | ttt | gcc | ttc | tgc | cgg   | gaa | tgt | aaa | gaa | gcg | tac | cat | 1220 |
| Leu | Gly | Cys | Gly | Phe | Ala | Phe | Cys | Arg   | Glu | Cys | Lys | Glu | Ala | Tyr | His |      |
|     |     | 360 |     |     |     |     | 365 |       |     |     |     | 370 |     |     |     |      |
| gaa | ggg | gag | tgc | agt | gcc | gta | ttt | gaa   | gcc | tca | gga | aca | act | act | cag | 1268 |
| Glu | Gly | Glu | Cys | Ser | Ala | Val | Phe | Glu   | Ala | Ser | Gly | Thr | Thr | Thr | Gln |      |
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| gcc | tac | aga | gtc | gat | gaa | aga | gcc | gcc   | gag | cag | gct | cgt | tgg | gaa | gca | 1316 |
| Ala | Tyr | Arg | Val | Asp | Glu | Arg | Ala | Ala   | Glu | Gln | Ala | Arg | Trp | Glu | Ala |      |
| 390 |     |     |     |     | 395 |     |     |       |     | 400 |     |     |     |     | 405 |      |
| gcc | tcc | aaa | gaa | acc | atc | aag | aaa | acc   | acc | aag | ccc | tgt | ccc | cgc | tgc | 1364 |
| Ala | Ser | Lys | Glu | Thr | Ile | Lys | Lys | Thr   | Thr | Lys | Pro | Cys | Pro | Arg | Cys |      |
|     |     |     |     | 410 |     |     |     |       | 415 |     |     |     |     | 420 |     |      |
| cat | gta | cca | gtg | gaa | aaa | aat | gga | ggc   | tgc | atg | cac | atg | aag | tgt | ccg | 1412 |
| His | Val | Pro | Val | Glu | Lys | Asn | Gly | Gly   | Cys | Met | His | Met | Lys | Cys | Pro |      |
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| Arg | Gln | Gly | Val | Pro | Ala | Asp | Gln | Leu | Arg | Val | Ile | Phe | Ala | Gly | Lys | 35  | 40  | 45  |     |
| Glu | Leu | Arg | Asn | Asp | Trp | Thr | Val | Gln | Asn | Cys | Asp | Leu | Asp | Gln | Gln | 50  | 55  | 60  |     |
| Ser | Ile | Val | His | Ile | Val | Gln | Arg | Pro | Trp | Arg | Lys | Gly | Gln | Glu | Met | 65  | 70  | 75  | 80  |
| Asn | Ala | Thr | Gly | Gly | Asp | Asp | Pro | Arg | Asn | Ala | Ala | Gly | Gly | Cys | Glu | 85  | 90  | 95  |     |
| Arg | Glu | Pro | Gln | Ser | Leu | Thr | Arg | Val | Asp | Leu | Ser | Ser | Ser | Val | Leu | 100 | 105 | 110 |     |
| Pro | Gly | Asp | Ser | Val | Gly | Leu | Ala | Val | Ile | Leu | His | Thr | Asp | Ser | Arg | 115 | 120 | 125 |     |
| Lys | Asp | Ser | Pro | Pro | Ala | Gly | Ser | Pro | Ala | Gly | Arg | Ser | Ile | Tyr | Asn | 130 | 135 | 140 |     |
| Ser | Phe | Tyr | Val | Tyr | Cys | Lys | Gly | Pro | Cys | Gln | Arg | Val | Gln | Pro | Gly | 145 | 150 | 155 | 160 |
| Lys | Leu | Arg | Val | Gln | Cys | Ser | Thr | Cys | Arg | Gln | Ala | Thr | Leu | Thr | Leu | 165 | 170 | 175 |     |
| Thr | Gln | Gly | Pro | Ser | Cys | Trp | Asp | Asp | Val | Leu | Ile | Pro | Asn | Arg | Met | 180 | 185 | 190 |     |
| Ser | Gly | Glu | Cys | Gln | Ser | Pro | His | Cys | Pro | Gly | Thr | Ser | Ala | Glu | Phe | 195 | 200 | 205 |     |
| Phe | Phe | Lys | Cys | Gly | Ala | His | Pro | Thr | Ser | Asp | Lys | Glu | Thr | Pro | Val | 210 | 215 | 220 |     |
| Ala | Leu | His | Leu | Ile | Ala | Thr | Asn | Ser | Arg | Asn | Ile | Thr | Cys | Ile | Thr | 225 | 230 | 235 | 240 |
| Cys | Thr | Asp | Val | Arg | Ser | Pro | Val | Leu | Val | Phe | Gln | Cys | Asn | Ser | Arg | 245 | 250 | 255 |     |
| His | Val | Ile | Cys | Leu | Asp | Cys | Phe | His | Leu | Tyr | Cys | Val | Thr | Arg | Leu | 260 | 265 | 270 |     |

Asn Asp Arg Gln Phe Val His Asp Pro Gln Leu Gly Tyr Ser Leu Pro  
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Cys Val Ala Gly Cys Pro Asn Ser Leu Ile Lys Glu Leu His His Phe  
290 295 300

Arg Ile Leu Gly Glu Glu Gln Tyr Asn Arg Tyr Gln Gln Tyr Gly Ala  
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325 330 335

Cys Gly Ala Gly Leu Leu Pro Glu Pro Asp Gln Arg Lys Val Thr Cys  
340 345 350

Glu Gly Gly Asn Gly Leu Gly Cys Gly Phe Ala Phe Cys Arg Glu Cys  
355 360 365

Lys Glu Ala Tyr His Glu Gly Glu Cys Ser Ala Val Phe Glu Ala Ser  
370 375 380

Gly Thr Thr Thr Gln Ala Tyr Arg Val Asp Glu Arg Ala Ala Glu Gln  
385 390 395 400

Ala Arg Trp Glu Ala Ala Ser Lys Glu Thr Ile Lys Lys Thr Thr Lys  
405 410 415

Pro Cys Pro Arg Cys His Val Pro Val Glu Lys Asn Gly Gly Cys Met  
420 425 430

His Met Lys Cys Pro Gln Pro Gln Cys Arg Leu Glu Trp Cys Trp Asn  
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                               Met Ile Val Phe Val
                               1               5

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Arg Phe Asn Ser Ser His Gly Phe Pro Val Glu Val Asp Ser Asp Thr
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agc atc ttc cag ctc aag gag gtg gtt gct aag cga cag ggg gtt ccg      212
Ser Ile Phe Gln Leu Lys Glu Val Val Ala Lys Arg Gln Gly Val Pro
                25                30                35

gct gac cag ttg cgt gtg att ttc gca ggg aag gag ctg agg aat gac      260
Ala Asp Gln Leu Arg Val Ile Phe Ala Gly Lys Glu Leu Arg Asn Asp
                40                45                50

tgg act gtg cag aat tgt gac ctg gat cag cag agc att gtt cac att      308
Trp Thr Val Gln Asn Cys Asp Leu Asp Gln Gln Ser Ile Val His Ile
                55                60                65

gtg cag aga ccg tgg aga aaa ggt caa gaa atg aat gca act gga ggc      356
Val Gln Arg Pro Trp Arg Lys Gly Gln Glu Met Asn Ala Thr Gly Gly
                70                75                80                85

gac gac ccc aga aac gcg gcg gga ggc tgt gag cgg gag ccc cag agc      404
Asp Asp Pro Arg Asn Ala Ala Gly Gly Cys Glu Arg Glu Pro Gln Ser
                90                95                100

ttg act cgg gtg gac ctc agc agc tca gtc ctc cca gga gac tct gtg      452
Leu Thr Arg Val Asp Leu Ser Ser Ser Val Leu Pro Gly Asp Ser Val
                105                110                115

ggg ctg gct gtc att ctg cac act gac agc agg aag gac tca cca cca      500
Gly Leu Ala Val Ile Leu His Thr Asp Ser Arg Lys Asp Ser Pro Pro
                120                125                130

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Ala Gly Ser Pro Ala Gly Arg Ser Ile Tyr Asn Ser Phe Tyr Val Tyr
                135                140                145

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Cys Lys Gly Pro Cys Gln Arg Val Gln Pro Gly Lys Leu Arg Val Gln
                150                155                160                165

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Cys Ser Thr Cys Arg Gln Ala Thr Leu Thr Leu Thr Gln Glu Phe Phe
                170                175                180

ttt aaa tgt gga gca cac ccc acc tct gac aag gaa aca cca gta gct      692
Phe Lys Cys Gly Ala His Pro Thr Ser Asp Lys Glu Thr Pro Val Ala
                185                190                195

ttg cac ctg atc gca aca aat agt cgg aac atc act tgc att acg tgc      740
Leu His Leu Ile Ala Thr Asn Ser Arg Asn Ile Thr Cys Ile Thr Cys
                200                205                210

aca gac gtc agg agc ccc gtc ctg gtt ttc cag tgc aac tcc cgc cac      788
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| Val Ile Cys Leu Asp Cys Phe His Leu Tyr Cys Val Thr Arg Leu Asn    |     |     |      |
| 230  | 235 | 240 | 245  |
| gat cgg cag ttt gtt cac gac cct caa ctt ggc tac tcc ctg cct tgt    |     |     | 884  |
| Asp Arg Gln Phe Val His Asp Pro Gln Leu Gly Tyr Ser Leu Pro Cys    |     |     |      |
| 250  | 255 | 260 |      |
| gtg gct ggc tgt ccc aac tcc ttg att aaa gag ctc cat cac ttc agg    |     |     | 932  |
| Val Ala Gly Cys Pro Asn Ser Leu Ile Lys Glu Leu His His Phe Arg    |     |     |      |
| 265  | 270 | 275 |      |
| att ctg gga gaa gag cag tac aac cgg tac cag cag tat ggt gca gag    |     |     | 980  |
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| 280  | 285 | 290 |      |
| gag tgt gtc ctg cag atg ggg ggc gtg tta tgc ccc cgc cct ggc tgt    |     |     | 1028 |
| Glu Cys Val Leu Gln Met Gly Gly Val Leu Cys Pro Arg Pro Gly Cys    |     |     |      |
| 295  | 300 | 305 |      |
| gga gcg ggg ctg ctg ccg gag cct gac cag agg aaa gtc acc tgc gaa    |     |     | 1076 |
| Gly Ala Gly Leu Leu Pro Glu Pro Asp Gln Arg Lys Val Thr Cys Glu    |     |     |      |
| 310  | 315 | 320 | 325  |
| ggg ggc aat ggc ctg ggc tgt ggg ttt gcc ttc tgc cgg gaa tgt aaa    |     |     | 1124 |
| Gly Gly Asn Gly Leu Gly Cys Gly Phe Ala Phe Cys Arg Glu Cys Lys    |     |     |      |
| 330  | 335 | 340 |      |
| gaa gcg tac cat gaa ggg gag tgc agt gcc gta ttt gaa gcc tca gga    |     |     | 1172 |
| Glu Ala Tyr His Glu Gly Glu Cys Ser Ala Val Phe Glu Ala Ser Gly    |     |     |      |
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| aca act act cag gcc tac aga gtc gat gaa aga gcc gcc gag cag gct    |     |     | 1220 |
| Thr Thr Thr Gln Ala Tyr Arg Val Asp Glu Arg Ala Ala Glu Gln Ala    |     |     |      |
| 360  | 365 | 370 |      |
| cgt tgg gaa gca gcc tcc aaa gaa acc atc aag aaa acc acc aag ccc    |     |     | 1268 |
| Arg Trp Glu Ala Ala Ser Lys Glu Thr Ile Lys Lys Thr Thr Lys Pro    |     |     |      |
| 375  | 380 | 385 |      |
| tgt ccc cgc tgc cat gta cca gtg gaa aaa aat gga ggc tgc atg cac    |     |     | 1316 |
| Cys Pro Arg Cys His Val Pro Val Glu Lys Asn Gly Gly Cys Met His    |     |     |      |
| 390  | 395 | 400 | 405  |
| atg aag tgt ccg cag ccc cag tgc agg ctc gag tgg tgc tgg aac tgt    |     |     | 1364 |
| Met Lys Cys Pro Gln Pro Gln Cys Arg Leu Glu Trp Cys Trp Asn Cys    |     |     |      |
| 410  | 415 | 420 |      |
| ggc tgc gag tgg aac cgc gtc tgc atg ggg gac cac tgg ttc gac gtg    |     |     | 1412 |
| Gly Cys Glu Trp Asn Arg Val Cys Met Gly Asp His Trp Phe Asp Val    |     |     |      |
| 425  | 430 | 435 |      |
| tagccagggc ggccggggcgc cccatcgcca catcctgggg gagcataccc agtgtctacc |     |     | 1472 |
| ttcatttttct aattctcttt tcaaacacac acacacacgc gcgcgcgcgc acacacactc |     |     | 1532 |
| ttcaagtttt tttcaaagtc caactacagc caaattgcag aagaaactcc tggatccctt  |     |     | 1592 |
| tcactatgtc catgaaaaac agcagagtaa aattacagaa gaagctcctg aatccctttc  |     |     | 1652 |

agtttgtcca cacaagacag cagagccatc tgcgacacca ccaacaggcg ttctcagcct 1712  
ccggatgaca caaataccag agcacagatt caagtgcaat ccatgtatct gtatgggtca 1772  
ttctcacctg aattcgagac aggcagaatc agtagctgga gagagagttc tcacatttaa 1832  
tatcctgcct ttaccttca gtaaacacca tgaagatgcc attgacaagg tgtttctctg 1892  
taaaatgaac tgcagtgggt tctccaaact agattcatgg ctttaacagt aatgttctta 1952  
tttaaatttt cagaaagcat ctattcccaa agaaccacag gcaatagtca aaaacatttg 2012  
tttatcctta agaattccat ctatataaat cgcattaatc gaaataccaa ctatgtgtaa 2072  
atcaacttgt cacaaagtga gaaattatga aagttaattt gaatgttgaa tgtttgaatt 2132  
acaggaaga aatcaagtta atgtactttc attccctttc atgatttgca actttagaaa 2192  
gaaattgttt ttctgaaagt atcaccaaaa aatctatagt ttgattctga gtattcattt 2252  
tgcaacttgg agattttgct aatacatattg gctccactgt aaatttaata gataaagtgc 2312  
ctataaagga aacacgttta gaaatgattt caaaatgata ttcaatctta acaaaagtga 2372  
acattattaa atcagaatct ttaaagagga gcctttccag aactaccaa atgaagacac 2432  
gcccgactct ctccatcaga agggtttata cccctttggc acaccctctc tgtccaatct 2492  
gcaagtcca gggagctctg cataccaggg gttccccagg agagaccttc tcttaggaca 2552  
gtaaactcac tagaatattc cttatgttga catggattgg atttcagttc aatcaaactt 2612  
tcagcttttt ttctagccat tcacaacaca atcaaaagat taacaacact gcatgcggca 2672  
aaccgcatgc tcttaccac actacgcaga agagaaagta caaccactat cttttgttct 2732  
acctgtattg tctgacttct caggaagatc gtgaacataa ctgagggcat gagtctcact 2792  
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Met Ile Val Phe Val Arg Phe Asn Ser Ser His Gly Phe Pro Val Glu  
1 5 10 15

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|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Val | Asp | Ser | Asp | Thr | Ser | Ile | Phe | Gln | Leu | Lys | Glu | Val | Val | Ala | Lys | 20  | 25  | 30  |     |
| Arg | Gln | Gly | Val | Pro | Ala | Asp | Gln | Leu | Arg | Val | Ile | Phe | Ala | Gly | Lys | 35  | 40  | 45  |     |
| Glu | Leu | Arg | Asn | Asp | Trp | Thr | Val | Gln | Asn | Cys | Asp | Leu | Asp | Gln | Gln | 50  | 55  | 60  |     |
| Ser | Ile | Val | His | Ile | Val | Gln | Arg | Pro | Trp | Arg | Lys | Gly | Gln | Glu | Met | 65  | 70  | 75  | 80  |
| Asn | Ala | Thr | Gly | Gly | Asp | Asp | Pro | Arg | Asn | Ala | Ala | Gly | Gly | Cys | Glu | 85  | 90  | 95  |     |
| Arg | Glu | Pro | Gln | Ser | Leu | Thr | Arg | Val | Asp | Leu | Ser | Ser | Ser | Val | Leu | 100 | 105 | 110 |     |
| Pro | Gly | Asp | Ser | Val | Gly | Leu | Ala | Val | Ile | Leu | His | Thr | Asp | Ser | Arg | 115 | 120 | 125 |     |
| Lys | Asp | Ser | Pro | Pro | Ala | Gly | Ser | Pro | Ala | Gly | Arg | Ser | Ile | Tyr | Asn | 130 | 135 | 140 |     |
| Ser | Phe | Tyr | Val | Tyr | Cys | Lys | Gly | Pro | Cys | Gln | Arg | Val | Gln | Pro | Gly | 145 | 150 | 155 | 160 |
| Lys | Leu | Arg | Val | Gln | Cys | Ser | Thr | Cys | Arg | Gln | Ala | Thr | Leu | Thr | Leu | 165 | 170 | 175 |     |
| Thr | Gln | Glu | Phe | Phe | Phe | Lys | Cys | Gly | Ala | His | Pro | Thr | Ser | Asp | Lys | 180 | 185 | 190 |     |
| Glu | Thr | Pro | Val | Ala | Leu | His | Leu | Ile | Ala | Thr | Asn | Ser | Arg | Asn | Ile | 195 | 200 | 205 |     |
| Thr | Cys | Ile | Thr | Cys | Thr | Asp | Val | Arg | Ser | Pro | Val | Leu | Val | Phe | Gln | 210 | 215 | 220 |     |
| Cys | Asn | Ser | Arg | His | Val | Ile | Cys | Leu | Asp | Cys | Phe | His | Leu | Tyr | Cys | 225 | 230 | 235 | 240 |
| Val | Thr | Arg | Leu | Asn | Asp | Arg | Gln | Phe | Val | His | Asp | Pro | Gln | Leu | Gly | 245 | 250 | 255 |     |
| Tyr | Ser | Leu | Pro | Cys | Val | Ala | Gly | Cys | Pro | Asn | Ser | Leu | Ile | Lys | Glu |     |     |     |     |

|   |     |         |
|---|-----|---------|
| 260   | 265 | 270     |
| Leu His His Phe Arg Ile Leu Gly Glu Glu Gln Tyr Asn Arg Tyr Gln |     |         |
| 275   | 280 | 285     |
| Gln Tyr Gly Ala Glu Glu Cys Val Leu Gln Met Gly Gly Val Leu Cys |     |         |
| 290   | 295 | 300     |
| Pro Arg Pro Gly Cys Gly Ala Gly Leu Leu Pro Glu Pro Asp Gln Arg |     |         |
| 305   | 310 | 315 320 |
| Lys Val Thr Cys Glu Gly Gly Asn Gly Leu Gly Cys Gly Phe Ala Phe |     |         |
| 325   | 330 | 335     |
| Cys Arg Glu Cys Lys Glu Ala Tyr His Glu Gly Glu Cys Ser Ala Val |     |         |
| 340   | 345 | 350     |
| Phe Glu Ala Ser Gly Thr Thr Thr Gln Ala Tyr Arg Val Asp Glu Arg |     |         |
| 355   | 360 | 365     |
| Ala Ala Glu Gln Ala Arg Trp Glu Ala Ala Ser Lys Glu Thr Ile Lys |     |         |
| 370   | 375 | 380     |
| Lys Thr Thr Lys Pro Cys Pro Arg Cys His Val Pro Val Glu Lys Asn |     |         |
| 385   | 390 | 395 400 |
| Gly Gly Cys Met His Met Lys Cys Pro Gln Pro Gln Cys Arg Leu Glu |     |         |
| 405   | 410 | 415     |
| Trp Cys Trp Asn Cys Gly Cys Glu Trp Asn Arg Val Cys Met Gly Asp |     |         |
| 420   | 425 | 430     |
| His Trp Phe Asp Val   |     |         |
| 435   |     |         |

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21

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<211> 22

<212> DNA

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gaaggtccca tttttcggtt tc

22

<210> 7

<211> 15

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<223> Xaa may be a combination of any 2 amino acids

<220>

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<222> (4) .. (4)

<223> Xaa may be a combination of any 9 amino acids

<220>

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<222> (6) .. (6)



<223> Xaa may be any amino acid

<220>

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<222> (8) .. (8)

<223> Xaa may be any combination of 2 amino acids

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<222> (10) .. (10)

<223> Xaa may be any combination of 4 amino acids

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<222> (12) .. (12)

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<222> (14) .. (14)

<223> Xaa may be any combination of 2 amino acids

<400> 7

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|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Cys | Xaa | Cys | Xaa | Cys | Xaa | His | Xaa | Cys | Xaa | Cys | Xaa | Cys | Xaa | Cys |
| 1   |     |     |     | 5   |     |     |     | 10  |     |     |     | 15  |     |     |

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<221> REPEAT

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<223> Xaa may be repeated 9 to 399 times

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<222> (6) .. (6)

<223> Xaa may be any amino acid

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<222> (8) .. (8)

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<223> Xaa may be repeated from 4 to 48 times

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<400> 8

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|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| Cys | Xaa | Cys | Xaa | Cys | Xaa | His | Xaa | Cys | Xaa | Cys | Xaa | Cys | Xaa | Cys |
| 1   |     |     |     | 5   |     |     |     | 10  |     |     |     | 15  |     |     |

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<400> 9  
accatgatag gtacgtgggt

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<210> 11

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<210> 21

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22

<210> 34

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19

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24

<210> 36

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gcggcgcaga gaggctgtac

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atgttgctat caccatttaa ggg

23

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tgatagtcac aactgtgtgt aag

23

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actgtctcat tagcgtctat ctt

23

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<210> 60

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<400> 60

Met Gln Ile Phe Val Lys Thr Leu Thr Gly Lys Thr Ile Thr Leu Glu



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                    20                      25                      30  
Lys Glu Gly Ile Pro Pro Asp Gln Gln Arg Leu Ile Phe Ala Gly Lys  
                    35                      40                      45  
Gln Leu Glu Asp Gly Arg Thr Leu Ser Asp Tyr Asn Ile Gln Lys Glu  
                    50                      55                      60  
Ser Thr Leu His Leu Val Leu Arg Leu Arg Gly Gly  
65                      70                      75

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<211> 76

<212> PRT

<213> Saccharomyces sp.

<400> 61

Met Gln Ile Phe Val Lys Thr Leu Thr Gly Lys Thr Ile Thr Leu Glu  
1                      5                      10                      15  
Val Glu Ser Ser Asp Thr Ile Asp Asn Val Lys Ser Lys Ile Gln Asp  
                    20                      25                      30  
Lys Glu Gly Ile Pro Pro Asp Gln Gln Arg Leu Ile Phe Ala Gly Lys  
                    35                      40                      45  
Gln Leu Glu Asp Gly Arg Thr Leu Ser Asp Tyr Asn Ile Gln Lys Glu  
                    50                      55                      60  
Ser Thr Leu His Leu Val Leu Arg Leu Arg Gly Gly  
65                      70                      75

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<211> 76

<212> PRT

<213> Glycine sp.

<400> 62

Met Gln Ile Phe Val Lys Thr Leu Thr Gly Lys Thr Ile Thr Leu Glu  
1 5 10 15

Val Glu Ser Ser Asp Thr Ile Asp Asn Val Lys Ala Lys Ile Gln Asp  
20 25 30

Lys Glu Gly Ile Pro Pro Asp Gln Gln Arg Leu Ile Phe Ala Gly Lys  
35 40 45

Gln Leu Glu Asp Gly Arg Thr Leu Ala Asp Tyr Asn Ile Gln Lys Glu  
50 55 60

Ser Thr Leu His Leu Val Leu Arg Leu Arg Gly Gly  
65 70 75

<210> 63

<211> 33

<212> DNA

<213> Homo sapiens

<400> 63

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33

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<212> DNA

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tcccaaagggt ccattcttgct gggatgatgt ttttaattcca

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<211> 38

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1 5 10

38

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<211> 10

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<213> Homo sapiens

<400> 67

Gly Pro Ser Cys Trp Asp Asp Val Leu Ile  
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Val His Leu Ala Gly Met Met Phe  
1 5

37

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<400> 69

Val His Leu Ala Gly Met Met Phe  
1 5

<210> 70

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<212> DNA

<213> Homo sapiens

<400> 70  
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21